

SEQUENCE LISTING

<110> Stern, David
Yan, Shi Du

<120> A MODEL OF ALZHEIMER'S-TYPE PATHOLOGY DOUBLE TRANSGENIC
MICE: ABAD AND APP (MUTANT)

<130> 0575/62176

<140> Not Yet Known

<141> 2000-08-14

<160> 7

<170> PatentIn Ver. 2.1

<210> 1

<211> 261

<212> PRT

<213> RAT

<400> 1

Met Ala Ala Ala Val Arg Ser Val Lys Gly Leu Val Ala Val Ile Thr
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Gly Gly Ala Ser Gly Leu Gly Leu Ser Thr Ala Lys Arg Leu Val Gly
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Gln Gly Ala Thr Ala Val Leu Leu Asp Val Pro Asn Ser Glu Gly Glu
35 40 45

Thr Glu Ala Lys Lys Leu Gly Gly Asn Cys Ile Phe Ala Pro Ala Asn
50 55 60

Val Thr Ser Glu Lys Glu Val Gln Ala Ala Leu Thr Leu Ala Lys Glu
65 70 75 80

Lys Phe Gly Arg Ile Asp Val Ala Val Asn Cys Ala Gly Ile Ala Val
85 90 95

Ala Ile Lys Thr Tyr His Glu Lys Lys Asn Gln Val His Thr Leu Glu
100 105 110

Asp Phe Gln Arg Val Ile Asn Val Asn Leu Ile Gly Thr Phe Asn Val
115 120 125

Ile Arg Leu Val Ala Gly Val Met Gly Gln Asn Glu Pro Asp Gln Gly

| | | |
|---|-----|---------|
| 130 | 135 | 140 |
| Gly Gln Arg Gly Val Ile Ile Asn Thr Ala Ser Val Ala Ala Phe Glu | | |
| 145 | 150 | 155 160 |
| Gly Gln Val Gly Gln Ala Ala Tyr Ser Ala Ser Lys Gly Gly Ile Val | | |
| | 165 | 170 175 |
| Gly Met Thr Leu Pro Ile Ala Arg Asp Leu Ala Pro Ile Gly Ile Arg | | |
| | 180 | 185 190 |
| Val Val Thr Ile Ala Pro Gly Leu Phe Ala Thr Pro Leu Leu Thr Thr | | |
| | 195 | 200 205 |
| Leu Pro Asp Lys Val Arg Asn Phe Leu Ala Ser Gln Val Pro Phe Pro | | |
| | 210 | 215 220 |
| Ser Arg Leu Gly Asp Pro Ala Glu Tyr Ala His Leu Val Gln Met Val | | |
| 225 | 230 | 235 240 |
| Ile Glu Asn Pro Phe Leu Asn Gly Glu Val Ile Arg Leu Asp Gly Ala | | |
| | 245 | 250 255 |
| Ile Arg Met Gln Pro | | |
| | 260 | |

<210> 2
 <211> 261
 <212> PRT
 <213> Human

<400> 2

| | | |
|---|----|-------|
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| Gly Gly Ala Ser Gly Leu Gly Leu Ala Thr Ala Glu Arg Leu Val Gly | | |
| | 20 | 25 30 |
| Gln Gly Ala Ser Ala Val Leu Leu Asp Leu Pro Asn Ser Gly Gly Glu | | |
| | 35 | 40 45 |
| Ala Gln Ala Lys Lys Leu Gly Asn Asn Cys Val Phe Ala Pro Ala Asp | | |
| 50 | 55 | 60 |
| Val Thr Ser Glu Lys Asp Val Gln Thr Ala Leu Ala Leu Ala Lys Gly | | |
| 65 | 70 | 75 80 |

Lys Phe Gly Arg Val Asp Val Ala Val Asn Cys Ala Gly Ile Ala Val
 85 90 95
 Ala Ser Lys Thr Tyr Asn Leu Lys Lys Gly Gln Thr His Thr Leu Glu
 100 105 110
 Asp Phe Gln Arg Val Leu Asp Val Asn Leu Met Gly Thr Phe Asn Val
 115 120 125
 Ile Arg Leu Val Ala Gly Glu Met Gly Gln Asn Glu Pro Asp Gln Gly
 130 135 140
 Gly Gln Arg Gly Val Ile Ile Asn Thr Ala Ser Val Ala Ala Phe Glu
 145 150 155 160
 Gly Gln Val Gly Gln Ala Ala Tyr Ser Ala Ser Lys Gly Gly Ile Val
 165 170 175
 Gly Met Thr Leu Pro Ile Ala Arg Asp Leu Ala Pro Ile Gly Ile Arg
 180 185 190
 Val Met Thr Ile Ala Pro Gly Leu Phe Gly Thr Pro Leu Leu Thr Ser
 195 200 205
 Leu Pro Glu Lys Val Cys Asn Phe Leu Ala Ser Gln Val Pro Phe Pro
 210 215 220
 Ser Arg Leu Gly Asp Pro Ala Glu Tyr Ala His Leu Val Gln Ala Ile
 225 230 235 240
 Ile Glu Asn Pro Phe Leu Asn Gly Glu Val Ile Arg Leu Asp Gly Ala
 245 250 255
 Ile Arg Met Gln Pro
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 <211> 973
 <212> DNA
 <213> Human

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 ctgcccttcc ttccctggg gtactactct ccagtcttgg gaggaagccc agtagccatt 900
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 aaaaaaaaaaaa aaa 973

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<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: PCR Primers

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<210> 5

<211> 19

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: PCR Primers

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19

<210> 6

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR Primers

<400> 6

gacaagtatc tcgagacacc tggggatgag

30

<210> 7

<211> 29

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
PCR Primer

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aaagaacttg taggttggat ttcgtacc

29